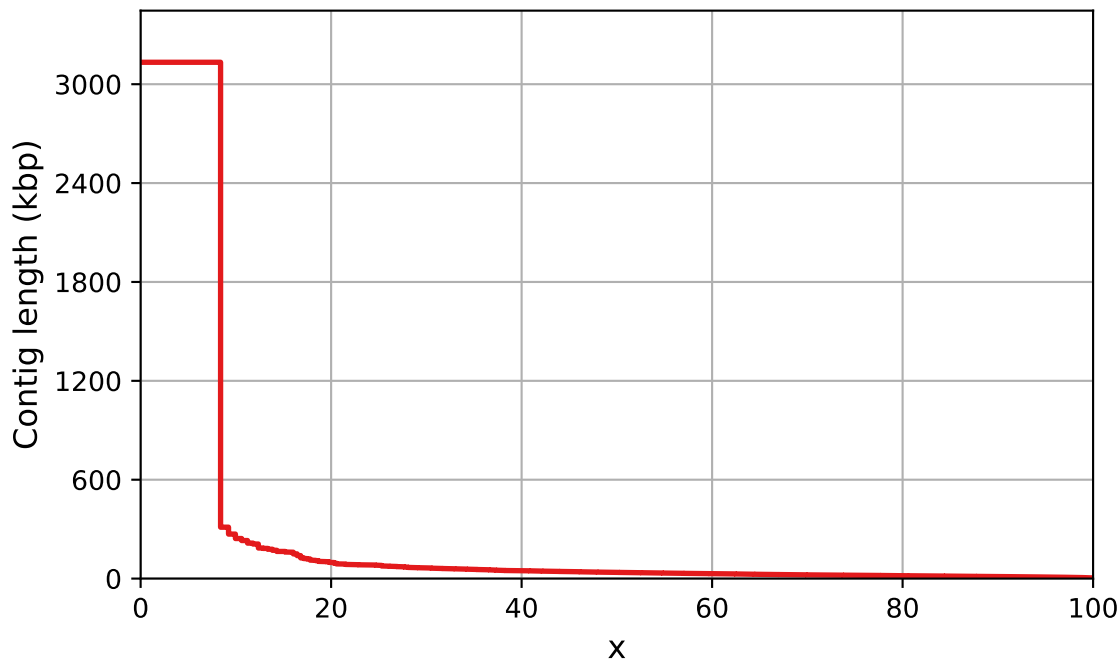


Report

	genome.nextpolish
# contigs (>= 0 bp)	1369
# contigs (>= 1000 bp)	1367
# contigs (>= 5000 bp)	1334
# contigs (>= 10000 bp)	1070
# contigs (>= 25000 bp)	424
# contigs (>= 50000 bp)	128
Total length (>= 0 bp)	37324549
Total length (>= 1000 bp)	37323051
Total length (>= 5000 bp)	37189619
Total length (>= 10000 bp)	35181904
Total length (>= 25000 bp)	24609226
Total length (>= 50000 bp)	14174980
# contigs	1369
Largest contig	3133102
Total length	37324549
GC (%)	52.06
N50	37673
N90	12307
auN	311994.1
L50	232
L90	928
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	91 + 9 part

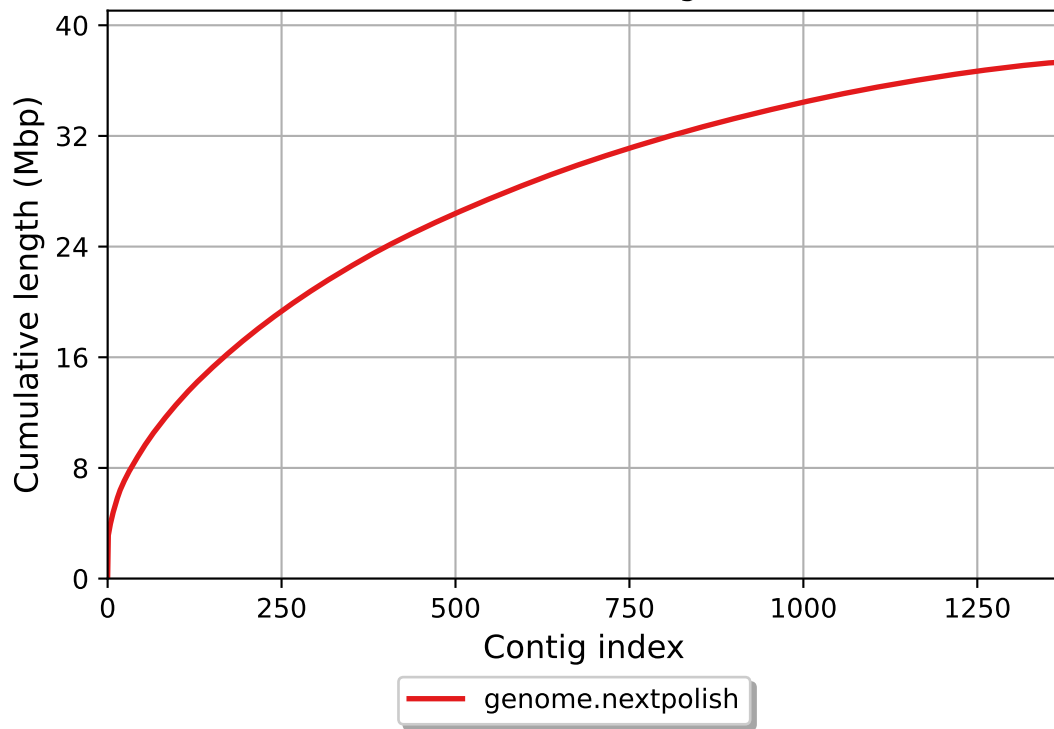
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

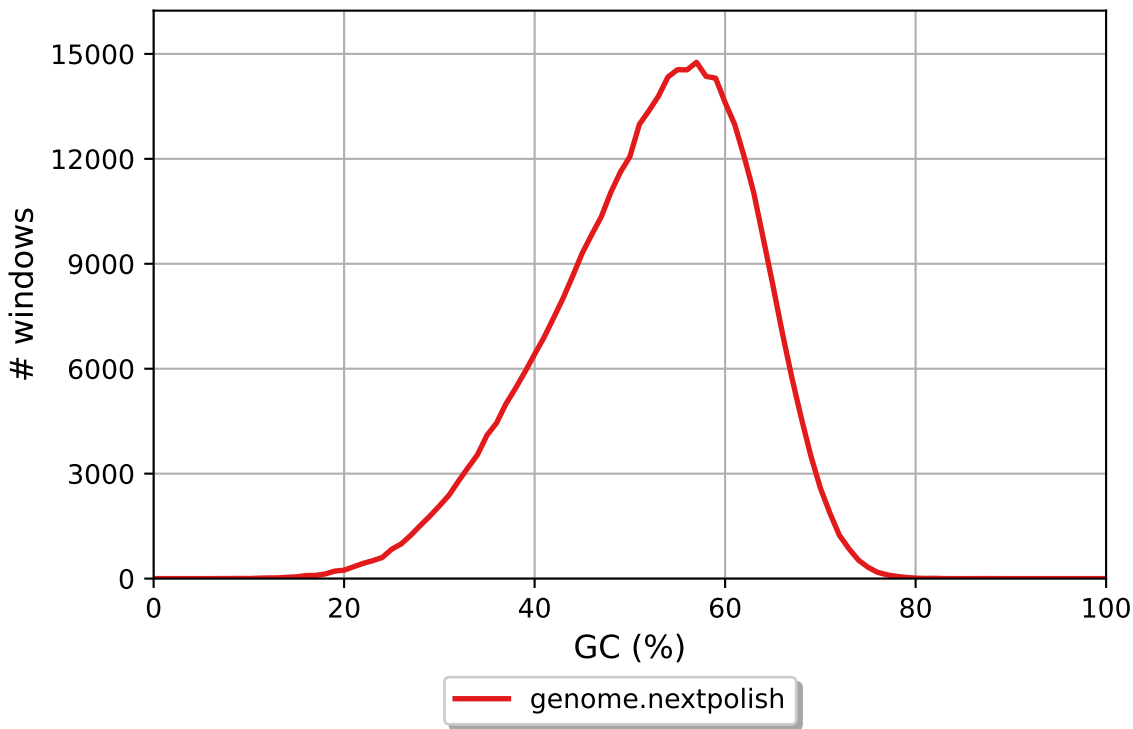


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

